

## INFLUENCE OF A GENETICALLY MODIFIED ENDOPHYTIC BACTERIUM ON COMPOSITION AND DECOMPOSITION OF CORN RESIDUE

C. F. TESTER

Soil-Microbial Systems Laboratory, ARS, USDA, Beltsville, MD 20705, U.S.A.

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**Summary**—I have determined whether the presence of an endophytic bacterium alters the chemical properties of inoculated plants and compared decomposition of colonized residues with control residues in soil. Greenhouse-grown corn (*Zea mays* L.) plants (18 days old) were inoculated with either the endophytic bacteria *Clavibacter xyli* subsp. *cynodontis* (MDE1) or a genetically engineered construction of *C. xyli* subsp. *cynodontis* which was transformed by inserting a gene encoding for production of a  $\delta$ -endotoxin from *Bacillus thuringiensis* subsp. *kurstaki* (MDR1.3). Leaves and stalks were harvested at maturity and various chemical and physical properties analyzed. In addition, leaf and stalk residues were incorporated into soil to determine the extent of their decomposition in a 63 day laboratory incubation study. Leaf residues from plants inoculated with MDR1.3 retained significantly more water than did those inoculated with buffer or MDE1. Leaves of inoculated plants contained significantly more C than did stalks of inoculated plants. Soluble proteins were the major contributors to the C content of leaves, whereas, soluble carbohydrates were predominant in the stalks. Inoculation with either MDR1.3 or MDE1 appeared to increase the quantity of N present in all plant parts. During early stages, leaf residues contained more readily decomposable substrates than did stalk residues. Leaves from plants inoculated with buffer decomposed significantly more after 63 days than those from plants inoculated with MDE1 or MDR1.3. In contrast, corn stalks inoculated with MDR1.3 decomposed significantly more than those inoculated with buffer or MDE1. Over the 63 day incubation ca 33% of the corn residues were decomposed. Although the presence of the endophytes, MDE1 and MDR1.3, during plant growth had some influence on chemical and physical properties of the residues, the extent of residue decomposition in soil ( $\text{CO}_2$  evolved in 63 days) was not substantially different.

### INTRODUCTION

The bacterium, *Clavibacter xyli* subsp. *cynodontis* (Davis *et al.*, 1984) is commonly found in nature where it colonizes the vascular system of bermudagrass (*Cynodon dactylon* L.) (Liao and Chen, 1981; Davis and Augustin, 1984). This microorganism also colonizes several important crop species including corn (*Zea mays* L.) (Kostka *et al.*, 1988a). A novel delivery system which utilizes naturally occurring plant endophytic bacteria was developed to confer specific insect resistance to plants (Fahey, 1988). *C. xyli* subsp. *cynodontis* has been genetically altered to express the chromosomally inserted  $\delta$ -endotoxin gene from *Bacillus thuringiensis* subsp. *kurstaki* and antibiotic resistance markers (Turnover *et al.*, 1991). The recombinant strain MDR1.3 is a prototype version of a microbial pesticide that was tested in corn for control of the European corn borer (*Ostrinia nubilalis*) (Kostka *et al.*, 1988b). MDR1.3 systemically colonizes the xylem in stems, leaves and husks of inoculated corn plants and can reach populations in excess of  $10^7$  CFU  $\text{g}^{-1}$  plant tissue (Reeser and Kostka, 1988). The endophyte does not survive naturally outside the host plant (Tomasino *et al.*, 1988; Turner and Tomasino, 1988; Davis, 1989), does not persist beyond 18 days in green corn residues incor-

porated into soil (Turner and Tomasino, 1988) and is not seed transmitted (Reeser and Sommerfeld, 1988).

There are no reports specifically relating physiological interactions of *C. xyli* subsp. *cynodontis* with any annual host plant. Are non-pathological effects of this endophytic bacterium on a non-host plant expressed in altered transformations and partitioning of plant biomass components? Moreover, no studies have been conducted to determine whether decomposition of mature, colonized plant residues in soil is altered. The objectives of the present study were to investigate the effect of *C. xyli* subsp. *cynodontis* and the recombinant construction on chemical and physical properties that relate to decomposability of corn residues and to compare decomposition of corn residues grown with or without *C. xyli* subsp. *cynodontis* and the recombinant construction.

### MATERIALS AND METHODS

#### *Plant culture and inoculation*

Corn (*Zea mays* L. cv. DeKalb T1100) was grown in 4 litre pots containing Terralite Metro-mix 350 (Grace & Co., Cambridge, Mass.) in a greenhouse at Crop Genetics International, Hanover, Md. Nutrients, including trace elements, were supplied twice

weekly by applications of Nutriculture 20-20-20 (Plant Marvel Laboratories, Chicago Heights, Ill.). Iron in the form of Sequesterene 330 Fe (Ciba-Geigy, Greensboro, N.C.) was added at 1.2 g plant<sup>-1</sup> 2 weeks after planting. Eighteen days after seeding, 10 plants were inoculated with 3  $\mu$ l of  $2.83 \times 10^9$  cfu ml<sup>-1</sup> *C. xyli* subsp. *cynodontis* (MDE1) (designated as "wild-type" hereinafter) suspended in sterile phosphate buffered saline solution (PBS) [10 mM Potassium phosphate (pH 7) and 210 mM NaCl], and another 10 were inoculated with 3  $\mu$ l of  $1.43 \times 10^{10}$  CFU ml<sup>-1</sup> *C. xyli* subsp. *cynodontis* strain MDR1.3 (designated as "recombinant" hereinafter) which contained the  $\delta$ -endotoxin gene from *B. thuringiensis* subsp. *kurstaki* (Fahey, 1988). Ten control plants were inoculated with 3  $\mu$ l of sterile PBS containing no cells. Both stalk and leaf tissue were inoculated by needle puncture (Vidaver, 1977; Turner *et al.*, 1991). Colonization by both wild-type and recombinant bacteria was confirmed for each plant 15 days after inoculation to ensure that only colonized plants were used. Stem and leaf midrib sections were cut, surface sterilized in 10% sodium hypochlorite solution for 1 min, rinsed with sterile water, dipped into 95% ethanol and flamed. The end of each section was cut back about 5 mm, squeezed and touched onto GCab (selective for wild-type bacteria) and GCanti (selective for both wild-type and recombinant bacteria) media (Anderson and Flaherty, 1988; Turner *et al.*, 1991). Plates containing the imprints were incubated at 27°C for 10 days and then examined microscopically for the presence of endophyte.

The corn was succulent and forming grain at 77 days when 6 plants of each treatment were harvested. Plants were separated into leaves and stalk and dried at 70°C for 9 days before they were ground to <2 mm with a Wiley mill. Oven drying effectively rendered the endophyte non-viable. Ground plant parts were stored at 4°C until analyzed or subsequently used in the laboratory incubation study. The experimental treatments were repeated twice in field studies and twice in the greenhouse. Plant residues from one greenhouse study were used for the specific analyses described in this paper.

#### Plant residue analyses

For the analyses of soluble carbohydrates and proteins, duplicate 1 g samples of the 6 replicates of each treatment were homogenized in 15 ml of 80% ethanol with a Polytron PT 10/35 homogenizer (Brinkmann Instruments, Westbury, N.Y.) for 30 s. The homogenates were kept at 60°C for 15 min followed by centrifugation at 17,000 g for 15 min (Sorvall SS34 rotor). The pellets were re-extracted with 10 ml of 80% ethanol and with 8 ml of 50% ethanol as described above. The supernatants were combined and the volume reduced to 3 ml by flash evaporation (40°C). After adjusting the final volume to 7 ml with deionized water, 60 mg of activated charcoal (6–14 mesh) was added and mixed before

incubation for 5 min at 25°C. Charcoal was removed by centrifugation at 10,000 g for 10 min and total soluble carbohydrates (SC) were determined by the anthrone reaction (Ashwell, 1957). Individual sugars were determined by gas-liquid chromatography (Li *et al.*, 1988).

Starch and soluble protein were determined by homogenizing the pellets from the ethanol extractions in 10 ml 0.5 N NaOH for 30 s with a Polytron and incubating at 22°C for 15 min before neutralization with 10 ml of 0.5 N acetic acid. The samples were clarified at 10,000 g for 10 min, and 1 ml of the supernatant was removed and mixed with 1 ml of amyloglucosidase reagent [1 mg ml<sup>-1</sup> amyloglucosidase from *Rhizopus* (Sigma Chemical Co., St Louis, Mo) in 0.1 M sodium acetate buffer, pH 4.8]. The mixture was kept for 1 h at 30°C before 30 mg of activated charcoal was added and further incubated at 22°C for 15 min. Charcoal was removed before a 1 ml sample was assayed for glucose. A second sample (2 ml) of the neutralized homogenate was treated with charcoal and 1 ml assayed for glucose. Glucose was determined by the glucose oxidase test (Fleming and Pegler, 1963). Starch glucose was calculated from the difference in glucose content with or without starch digestion by amyloglucosidase. Soluble protein in the neutralized homogenates was assayed by the Bradford method (1976). Non-soluble C components were determined by subtracting the sum of the soluble carbohydrates (42% C) and protein (50% C) from total carbon.

#### Residue decomposition in soil

The soil used for the decomposition study was an Evesboro loamy sand subsoil (mesic, coated Typic Quartzipsamment, 97% sand) collected from a fescue meadow at the Beltsville Agriculture Research Center. This Upland Coastal plain soil has a pH of 4 and low amounts of total C, N, P and S, and has been used extensively in studies evaluating residue decomposition in soil (Tester *et al.*, 1977; Tester and Parr, 1983; Tester, 1988). The soil was air-dried and sieved ( $\leq 2$  mm). Lime (3.5 g of CaO: 5 kg soil) was added and mixed thoroughly to yield a pH of 6.4 and stored at 4°C until used in the laboratory incubation study (Tester, 1988). Water content at 33 kPa tension was determined by the pressure-plate method of Klute (1986) for soil, plant residues and 5% (w/w) mixtures of plant residues and soil.

5 g of the appropriate ground plant parts were mixed with 95 g of soil and placed into individual incubation flasks (6 cm dia  $\times$  12 cm high). The mixtures were adjusted to 33 kPa moisture tension, covered with plastic wrap and equilibrated for 3 days at 4°C before initiating the incubation study. Three replicates of each treatment were placed at random on a manifold system controlled at 25°C. The incoming air stream was passed through 4 N NaOH and then through distilled water to maintain the 33 kPa moisture tension (Tester and Parr, 1983). Moisture

Total C and N of the soil and plant residues were determined by dry combustion using an Erba automated analyzer (Carlo Erba Strumentazione, Milan, Italy). Data were analyzed by ANOVA and GLM (MEANS statement with LSD mean separation options) (SAS Institute, 1987). First-order decomposition rate constants were determined from the integrated rate equation expressed as

**where:**

$t$  = time (days).

Inoculated plants grew and matured at the same rate as did control plants and plant dry mass yields were not significantly different for all three treatments when harvested (Table 1) (no changes in dry mass yields due to inoculation were observed in either the field or greenhouse grown corn plants).

Total C, total N and water retention of corn plant residues are shown in Table 1. Ground leaves from plants inoculated with recombinant cells retained significantly more water than did leaves from either the control or wild-type inoculated plants. Ground stalks from control plants retained significantly more water than those from plants inoculated with either recombinant or wild-type cells. Leaf tissue from all treatments retained significantly more water than did the corresponding stalk tissue, probably the result of differences in chemical composition of the specific plant parts (Table 2). Soluble protein accounted for *ca* 22% of the total C found in leaf tissue, whereas, soluble carbohydrates represented *ca* 20% of the total C of stalk tissue. Collins *et al.* (1990) reported that wheat stems contained more lignin and nonstructural carbohydrates than did wheat leaves, which had higher soluble C. In this study, total C in corn residue probably reflected selective partitioning in plant parts from the different treatments. Even though not significant ( $P \leq 0.05$ ), leaves from wild-type and recombinant inoculated plants contained more C than did leaves from the control plants. In contrast, stalks from both wild-type and recombinant inoculated plants contained somewhat less C (again not significant at 0.05) than did stalks from the control plants. Note that leaves and stalks of control plants contained the same quantity of C. However, the quantity of C found in leaves of inoculated plants was significantly greater than that found in stalks of inoculated plants (Table 1). These total C values correspond to organic matter contents which determine the quantity of water retained by a substance (Klute, 1986). However, the distribution of organic compounds associated with leaves and stalks are different (Table 2) and thus result in differences in water retention.

Starch concentrations of corn residues were  $<0.5 \text{ mg g}^{-1}$  and did not significantly differ among treatments. Stalks contained approximately three

Material	Property				
	Dry wt	33 kPa†	Total C	Total N	C:N ratio
	g		g kg <sup>-1</sup> dry wt		
Control leaf	29b‡	2803b‡	482.4ab‡	26.25b‡	18.4
Wild-type leaf	28b	2799b	505.7a	28.35ab	17.8
Recombinant leaf	27b	2903a	502.1a	28.98a	17.3
Control stalk	80a	1695c	479.0ab	10.35c	44.9
Wild-type stalk	83a	1638d	464.8b	11.35c	41.0
Recombinant stalk	84a	1625d	464.7b	11.44c	40.6
1 SD (0.05)	4	45	23.4	2.20	4.4

†Means of six replications followed by the same letter are not significantly different at  $P \leq 0.05$ .

Table 2. Total carbon, soluble carbohydrates (SC, 42% C), soluble protein (50% C), nonsoluble C components (NSC) and ethanol soluble sugars of corn leaf and stalk samples (percentage of total carbon in parentheses)

Material	Total C	SC (%)	Protein (%)	NSC (%)†	Ethanol soluble sugars			
					Fructose	Glucose	Sucrose	Total
				mg g <sup>-1</sup> dry wt				
Control leaf	482.4ab†	82.4 (7.2)d†	216 (22.4)a*	654 (70.4)	3.2	2.4	33.1	38.7
Wild-type leaf	505.7a	72.5 (6.0)e	224 (22.1)a	699 (71.8)	3.3	2.3	33.2	38.8
Recombinant leaf	502.1a	79.4 (6.6)d	226 (22.5)a	684 (70.9)	3.2	2.9	37.1	43.3
Control stalk	479.0ab	233.4 (20.5)a	97 (10.1)b	692.7 (69)	37.8	35.7	41.7	115.2
Wild-type stalk	464.8b	218.0 (19.7)b	98 (10.5)b	675.4 (70)	35.6	29.5	44.2	109.3
Recombinant stalk	464.7b	208.0 (18.8)c	104 (11.2)b	677.7 (70)	36.4	31.2	45.3	112.9
LSD (0.05)	23.4	6.1	13					

\*Means of six replications followed by the same letter are not significantly different at  $P \leq 0.05$ .

†Nonsoluble C components were determined by subtracting the sum of soluble carbohydrates and protein from total C. Based on leaf and stalk structural components containing ca 52 and 48% C respectively.

times the quantity of soluble carbohydrates (SC-anthrone positive) as that detected in leaves (Table 2). The individual sugars (fructose, glucose, and sucrose) totaled ca 50% of the SC for each corn residue sample (Table 2). Sucrose accounted for 86% of the sugars found in leaf samples but only 40% of the sugars in stalk samples. The remainder was equally divided between fructose and glucose. No significant differences were detected for the individual sugars (fructose, glucose, and sucrose) found either in the leaves or stalks of individual treatments.

Total N analyses did not reflect the same partitioning trend as seen for total C. Leaves from recombinant inoculated plants contained significantly more N than did leaves from the control plants. Even though not always significant, inoculation with either wild-type or recombinant cells resulted in increased amounts of N in both leaves and stalks as compared to control plants. Stalks contained only 40% of the quantity of N found in leaves. This corresponds to the different types of organic compounds associated with different plant tissues (Table 2). Like total N, soluble protein content of the stalks was 45% of the protein content of leaves.

#### Residue decomposition in soil

C:N ratios of stalks differed radically from that of leaves, but only minor differences occurred among treatments (Table 1). Total C and N analyses and C:N ratios of plant residues alone are not definitive predictors of subsequent decomposition of residues in soil. The quantity, composition, and C:N ratios of the readily available organic constituents have been found to be more reliable indicators of rate and extent of decomposition of the material (Herman *et al.*, 1977; Reinersten *et al.*, 1984; Jawson and Elliott, 1986). Cumulative CO<sub>2</sub> evolution of the individual treatment residues is shown in Fig. 1 (presented for comparisons of rates of decomposition).

Decomposition rates of leaf substrates from all treatments were indistinguishable during the first 9 days (Table 3), but slight differences were noted

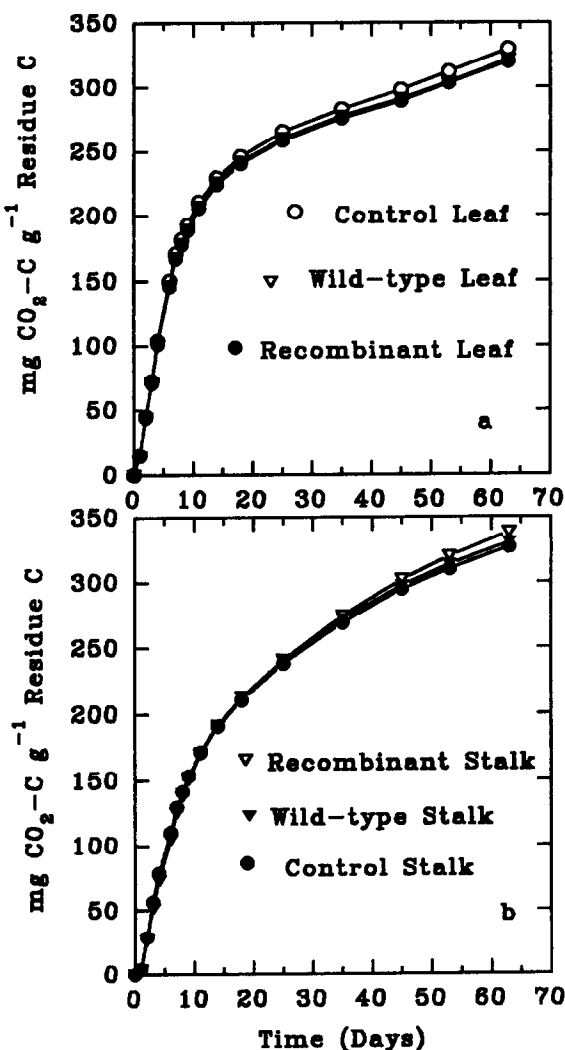


Fig. 1. Cumulative CO<sub>2</sub> evolved during the decomposition of 5 g of either corn leaf or corn stalk residue mixed with 95 g of Evesboro soil. (a) Leaf residue from control and endophyte colonized plants and (b) Stalk residue from control and endophyte colonized plants. See Table 1 for significance. Error bars are smaller than symbols for each data point.

Table 3. Decomposition stages and first-order rate constants† for corn residues mixed uniformly with Evesboro soil

Treatment Material	Stage I (0–9 days)	Stage II (10–35 days)	Stage III (40–63 days)
	rate constants (day <sup>-1</sup> )		
Control leaf	0.02559	0.00386*	0.00258*
Wild-type leaf	0.02515	0.00370	0.00248
Recombinant leaf	0.02494	0.00366	0.00247
Control stalk	0.02022	0.00512	0.00261
Wild-type stalk	0.02038	0.00523	0.00270
Recombinant stalk	0.02000	0.00545	0.00294*

\* $P \leq 0.05$ .  $F$ -test comparisons between treatments. All  $r^2 = 0.99$ .

†From the integrated rate equation  $C = C_0 e^{-kt}$ .

thereafter, with substrates in control leaves decomposing significantly more during the 63 days than those from inoculated leaves (Table 1). This either indicated that C-substrates present in inoculated leaves were less available than those present in control leaves or that more C fixation may have occurred. Reinertsen *et al.* (1984) reported that soluble and readily available C were directly related to the initial CO<sub>2</sub> evolved from wheat straw. Herman *et al.* (1977) showed that differences in CO<sub>2</sub> evolution from plant residues during decomposition was proportional to polysaccharide content and inversely proportional to lignin content and C:N ratios.

Decomposition rates of stalk substrates from all treatments were variable (Table 3), resulting in no difference in the quantity of CO<sub>2</sub>-C evolved for wild-type inoculated and control stalks. However, they decomposed significantly less than did the recombinant inoculated stalks (Table 1). Stalks from plants inoculated with recombinant contained the lowest C content and decomposed to the greatest extent in this study. The types of organic substances present in stalks from plants inoculated with recombinant may account for the observed differences in decomposition (trend toward more soluble protein but significantly less soluble carbohydrates, Table 2). The small difference in amount of SC present in stalks from plants inoculated with recombinant and the lack of difference in amounts of fructose, glucose and sucrose may indicate that some undetermined factors contributed to the observed differences in decomposition.

On average, corn residues in this study decomposed ca 33% during the 63 days incubation (Table 1), which is comparable to the decomposition of wheat straw in a similar incubation study (Tester, 1988). The differences observed in C and N concentrations and rate and extent of decomposition of corn residues in this controlled environment study are so small that they would likely be undetectable in field situations which are subject to wide ranges of environmental fluctuations.

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